

Fig. 1**SEQ ID NO: 1**

GGATTTCCGGGCTCCATGGCAAGATCCCTTCTCCTGCCCCTGCAGATCCTAC
TGCTATCCTTAGCCTTGGAACCTGCAGGAGAAGAAGCCCAGGGTGACAAGAT
TATTGATGGCGCCCCATGTGCAAGAGGCTCCCACCCATGGCAGGTGGCCCTG
CTCAGTGGCAATCAGCTCCACTGCGGAGGCGTCCTGGTCAATGAGCGCTGGG
TGCTCACTGCCGCCCCTGCAAGATGAATGAGTACACCGTGACCTGGGCAG
TGATACGCTGGGCGACAGGAGAGCTCAGAGGATCAAGGCCTCGAAGTCATTC
CGCCACCCCGGCTACTCCACACAGACCCATGTTAATGACCTCATGCTCGTGA
AGCTCAATAGCCAGGCCAGGCTGTCATCCATGGTGAAGAAAGTCAGGCTGCC
CTCCCGCTGCGAACCCCCCTGGAACCACCTGTACTGTCTCCGGCTGGGGCACT
ACCACGAGCCCAGATGTGACCTTTCCCTCTGACCTCATGTGCGTGGATGTCA
AGCTCATCTCCCCCAGGACTGCACGAAGGTTTACAAGGACTTACTGGAAAA
TTCCATGCTGTGCGCTGGCATCCCCGACTCCAAGAAAAACGCCTGCAATGGT
GACTCAGGGGGACCGTTGGTGTGCAGAGGTACCCTGCAAGGTCTGGTGTCTCT
GGGGAAC'TTCCCTTGCGGCCAACCCAATGACCCAGGAGTCTACACTCAAGT
GTGCAAGTTCACCAAGTGGATAAATGACACCATGAAAAAGCATCGCTAACGC
CACACTGAGTTAATTAATGCTGTGTGCTTCCAACAGAAAATGCACAGGAGTGAG
GACGCCGATGACCTATGAAGTCAAATTTGACTTTACCTTTCCTCAAAGATAT
ATTTAAACCTCATGCCCTGTTGATAAACCAATCAAATTGGTAAAGACCTAAA
ACCAAAACAAATAAAGAAACACAAAACCCCTCAA

Fig. 2**SEQ ID NO: 2**

MARSLLLPLQIILLLSLALETAGEEAAQGDKIIDGAPCARGSHPWQVALLSGNQ
LHCGGVLVNERWVLTAAHCKMNEYTVHLGSDTLGDRRAQRIKASKSFRHPGY
STQTHVNDLMLVKLNSQARLSSMVKKVRLPSRCEPPGTTCTVSGWGTTTSPD
VTFPSDLMCVDVKLISPQDCTKVYKDLLENSMLCAGIPDSKKNACNGDSGGP
LVCRGTLQGLVSWGTFPCGQPNDPGVYTQVCKFTKWINDTMKKHR

Fig. 3**SEQ ID NO: 3**

5' - GCATCCCCGACTCCAAGAA -3'

Fig. 4**SEQ ID NO: 4**

5' - CAGGGTACCTCTGCACACCAA -3'

Fig. 5**SEQ ID NO: 5**

5' - AACGCCTGCAATGGTGACTCAGGG -3'